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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2009; month=9; day=4; hr=15; min=0; sec=47; ms=158; ]

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Reviewer Comments:

<110> Teunis Bernard Herman Geijtenbeek; Yvette Van Kooyk

Please show each applicant on a separate line; show the <110> only once, at the beginning of the first applicant's line. Remove the semicolon.

<151> 2002/11/07

The above date response is invalid: replace with 2002-11-07

<210> 1

<211> 17

<212> DNA

<213> Artificial sequence

<223> Synthetic nucleotide

<400> 1

AGACTGCGT ACCAATTC

Please insert a mandatory "<220>" above the <223> line. <220> is mandatory whenever <221>, <222>, or <223> is shown. Please edit the <223> response: since more than 1 nucleotide is shown, use "Synthetic construct", instead. Please use lower-case letters for nucleotides; show the cumulative nucleotide total at the right margin of each line. Same errors in Sequence 2.

<210> 3

<211> 44

<212> PRT

<213> Artificial sequence

<223> Synthetic peptide

<400> 3

Trp Met Gly Leu Ser Asp Leu Asn Gln Glu Gly Thr Trp Gln Trp Val Asp Gly  
Ser

1 5 10 15

Pro Leu Leu Pro Ser Phe Lys Gln Tyr Trp Asn Arg Gly Glu Pro Asn Asn Val  
Gly

20 25 30 35

Glu Glu Asp Cys Ala Glu

40

Please insert a <220> above <223> line. Per Sequence Rules, a maximum  
of 16 amino acids per line. Same error in Sequence 4.

<210> 5

<211> 78

<212> DNA

<213> Artificial sequence

<223> Synthetic nucleotide

<400> 5

ATGTTCCAAC CCCTATTAGA CGCCTTCATA GAAAGCGCTT CCATTGAAAA AATGGCCTCT 60

AAATCTCCCC CCCCCTAA 78

Insert a <220> above <223>; replace the "<223> Synthetic nucleotide"  
with "<223> Synthetic construct". Use lower-case letters for  
nucleotides. Do not use TAB codes between the last nucleotide on a line  
and the cumulative nucleotide total. Same error in Sequence 6.

(end of Sequence 6)

AAATCTCCCC CCCCCTAAA AATC 84

1

Please remove the above numbers which appear at the end of the submitted file.

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Application No: 10533981

Version No: 1.0

Input Set:

Output Set:

Started: 2009-08-21 13:00:03.809

Finished: 2009-08-21 13:00:13.018

Elapsed: 0 hr(s) 0 min(s) 9 sec(s) 209 ms

Total Warnings: 201

Total Errors: 14

No. of SeqIDs Defined: 6

Actual SeqID Count: 6

Error code	Error Description
E 287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <141>
E 287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (1)
W 112	Upper case found in data; Found at position(0) SEQID(1)
W 112	Upper case found in data; Found at position(1) SEQID(1)
W 112	Upper case found in data; Found at position(2) SEQID(1)
W 112	Upper case found in data; Found at position(3) SEQID(1)
W 112	Upper case found in data; Found at position(4) SEQID(1)
W 112	Upper case found in data; Found at position(5) SEQID(1)
W 112	Upper case found in data; Found at position(6) SEQID(1)
W 112	Upper case found in data; Found at position(7) SEQID(1)
W 112	Upper case found in data; Found at position(8) SEQID(1)
W 112	Upper case found in data; Found at position(9) SEQID(1)
W 112	Upper case found in data; Found at position(10) SEQID(1)
W 112	Upper case found in data; Found at position(11) SEQID(1)
W 112	Upper case found in data; Found at position(12) SEQID(1)
W 112	Upper case found in data; Found at position(13) SEQID(1)
W 112	Upper case found in data; Found at position(14) SEQID(1)

**Input Set:**

**Output Set:**

**Started:** 2009-08-21 13:00:03.809  
**Finished:** 2009-08-21 13:00:13.018  
**Elapsed:** 0 hr(s) 0 min(s) 9 sec(s) 209 ms  
**Total Warnings:** 201  
**Total Errors:** 14  
**No. of SeqIDs Defined:** 6  
**Actual SeqID Count:** 6

Error code	Error Description
W 112	Upper case found in data; Found at position(15) SEQID(1)
W 112	Upper case found in data; Found at position(16) SEQID(1)
E 254	The total number of bases conflicts with running total Input: 0, Calculated : 17 SEQID(1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (2)
W 112	Upper case found in data; Found at position(0) SEQID(2)
W 112	Upper case found in data; Found at position(1) SEQID(2)
W 112	Upper case found in data; Found at position(2) SEQID(2) This error has occurred more than 20 times, will not be displayed
E 254	The total number of bases conflicts with running total Input: 0, Calculated : 16 SEQID(2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (6)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (6)

**Input Set:**

**Output Set:**

**Started:** 2009-08-21 13:00:03.809  
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**No. of SeqIDs Defined:** 6  
**Actual SeqID Count:** 6

Error code	Error Description
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (6)

# SEQUENCE LISTING

<110> Teunis Bernard Herman Geijtenbeek; Yvette Van Kooyk  
 <120> C-Type Lectin Binding Molecules, Identification and Uses Thereof  
 <130> 1943-2PCT/US  
 <140> 10533981  
 <141> 2009-08-21  
 <150> PCT/NL2003/000781  
 <151> 2002/11/07  
 <160> 6

<210> 1  
 <211> 17  
 <212> DNA  
 <213> Artificial sequence  
 <223> Synthetic nucleotide  
 <400> 1

AGACTGCGT ACCAATTC

<210> 2  
 <211> 16  
 <212> DNA  
 <213> Artificial sequence  
 <223> Synthetic nucleotide  
 <400> 2

GACGATGAGT CCTGAG

<210> 3  
 <211> 44  
 <212> PRT  
 <213> Artificial sequence  
 <223> Synthetic peptide  
 <400> 3

Trp Met Gly Leu Ser Asp Leu Asn Gln Glu Gly Thr Trp Gln Trp Val Asp Gly Ser  
 1 5 10 15

Pro Leu Leu Pro Ser Phe Lys Gln Tyr Trp Asn Arg Gly Glu Pro Asn Asn Val Gly





